Mevatonic acid Sterol-Specific Blosynthetic Pathway

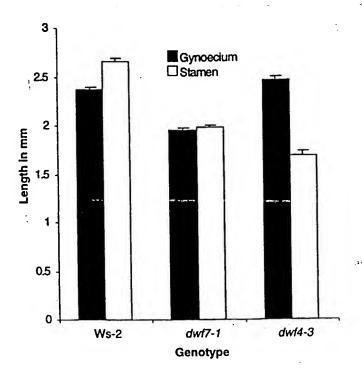


FIG. 2

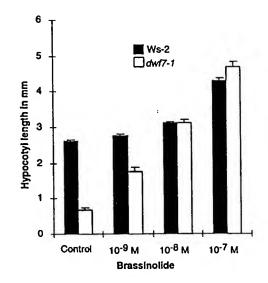


FIG. 3

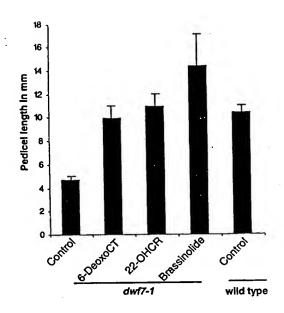
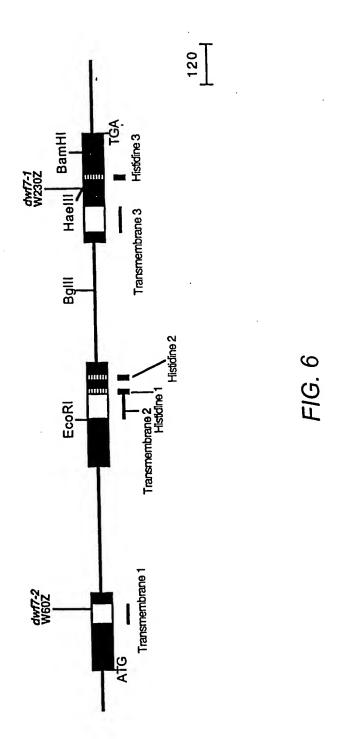


FIG. 4



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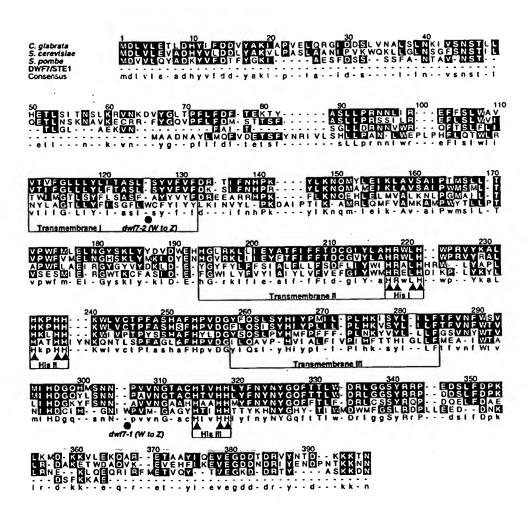


FIG. 7

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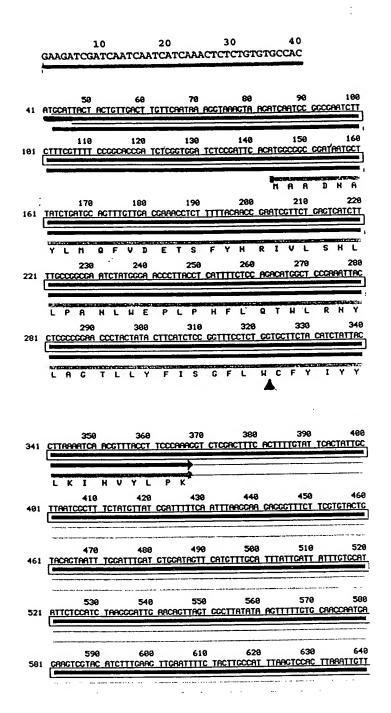


FIG. 8A

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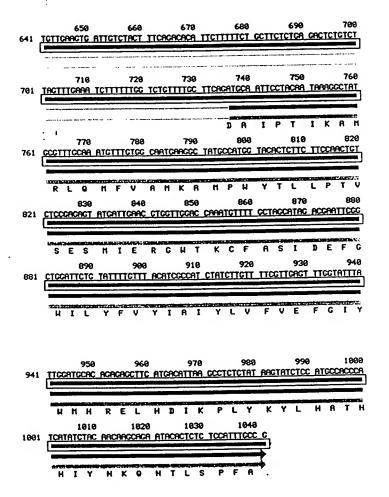


FIG. 8B

TETTETTING TTETTETARA CTRACTCTT TTCRCTTTCT ACRITECTAC CRITTACTIT CTTACCACAR ARCACTITET CACCACCTCC TIGIACTCCA TECRTTECTT ATCCATAGAG TRACCACAGA GCCTAGAATT ATATAGATGT TTACATCTTT CACCTCCRTT ACTTCACATA TGTCACAGAG ACTTCTCACT TAACCACAGT CTCTTTCTCT TCTCCTCTCC CACTCATTCC RARTCACCAC RACTTCTTTT ATCTACTTCC CTCCRCTCTA TCTTCCTTRA TCCRRCCRTC TCRCRTCTRA TATTACTTCT RACTTCCTTR CCRITTCRCC CACTACACCC CATACTTCRC CCTGTACCCC CCTTTTCTT 1450 1460 1470 1480 1490 1500 ATCTCATACC CCTCTTTATA CTCCCARTTC ATTTCACAAC TCATATACCT CTTTTGTTCA TOCCRACATO TOCCCACTAR TCCCTCCRCC RTRCCRTRCC ATRCRCCRCR CCRCRTRCRA CCRTRRCTAT CCTCRTTATA

FIG. 8C

IHH

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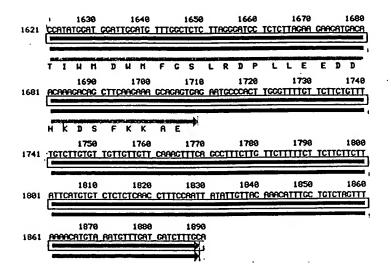


FIG. 8D

1	MAADNAYLMQ	FVDETSFYNR	IVLSHLLPAN	LWEPLPHFLQ	TWLRNYLAGI
51	LLYFISGFLW	CFYIYYLKIN	VYLPKDAIPT	IKAMRLQMFV	AMKAMPWYTI
101	LPTVSESMIE	RGWTKCFASI	DEFGWILYFV	YIAIYLVFVE	FGIYWMHREI
151	HDIKPLYKYL	HATHHIYNKQ	NTLSPFAGLA	FHPVDGILQA	VPHVIALFI
201	PIHFTTHIGL	LFMEAIWTAN	IHDCIHGNIW	PVMGAGYHTI	HHTTYKHNYO
251	HYTIWMDWMF	GSLRDPLLEE	DDNKDSFKKA	E	

FIG. 9

DWF 7 MUTANTS 50 10 30 GTTTGGTATTTATTGGATGCACAGAGAGCTTCATGACATTAAGCCTCTCTATAAGTATCT CAAACCATAAATAACCTACGTGTCTCTCGAAGTACTGTAATTCGGAGAGATATTCATAGA 110 90 70 CCATGCCACCCATCATATCTACAACAAGCAGAATACACTCTCTCCATTTGCCGGTAAGTG GGTACGGTGGGTAGTATAGATGTTGTTCGTCTTATGTGAGAGAGGTAAACGGCCATTCAC 170 150 130 TTTTCAGTTTGTTCTTTAGTTCTTGTAAAAGATTGGTAGCATTTAGTTTCTTACCAG AAAAGTCAAACAAGAAGAAATCAAGAACATTTTCTAACCATCGTAAATCAAAGAATGGTC 230 210 190 AAAAGACTTTGTCAGCAGCTGCTTGTACTCCAAATCACATTTTGCATTCCTTATCCATAA TTTTCTGAAACAGTCGTCGACGAACATGAGGTTTAGTGTAAAACGTAAGGAATAGGTATT 290 250 270 AGTAACCAGAAAGGCTAGAATTATATAAATGTCAGCTGCATTACTTCACATATGTCAGAG TCATTGGTCTTTCCGATCTTAATATATTTACAGTCGACGTAATGAAGTGTATACAGTCTC 350 330 310 AGACTTCTGACTTAACCAGAGTTTAGATCTTTGTGTTTCTCTTCTGGTCTCGGACTGATT TCTGAAGACTGAATTGGTCTCAAATCTAGAAACACAAAGAGAAGACCAGAGCCTGACTAA 410 370 390 GGAAATGACGAGAAGTTCTTTTATCTACTTCCCTGGAGTGTATCTTGGTTAATCCAAGGA CCTTTACTGCTCTTCAAGAAAATAGATGAAGGGACCTCACATAGAACCAATTAGGTTCCT 470 450 430 TGTGACATCTAAATATTACTTGTAACTTCCTTACGTTTTTGTTTACAGGGCTTGCATTCA ACACTGTAGATTTATAATGAACATTGAAGGAATGCAAAAACAAATGTCCCGAACGTAAGT 490 510 530 CCCAGTAGACGGGATACTTAAGGCTGTACCGCATGTGATAGCGCTGTTATAGTGCCAATT GGGTCATCTGCCCTATGAATTCCGACATGGCGTACACTATCGCGACAATATCACGGTTAA 590 570 550 CATTTCACAACTCATATAGGTCTTTTGTTCATGGAAGCGATATGGACGGCGAACATCCAT

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Matter No.: 11696-069002

Applicant(s): Sunghwa Choe et al.

FIG. 10A

GTAAAGTGTTGAGTATATCCAGAAAACAAGTACCTTCGCTATACCTGCCGCTTGTAGGTA

Matter No.: 11696-069002 Page 13 of 18 Applicant(s): Sunghwa Choe et al. **DWF 7 MUTANTS** GACTGCATCCATGGCAACATCTGGCCAGTAATGGGTGCAGGATACCATACGATACACCAC CTGACGTAGGTACCGTTGTAGACCGGTCATTACCCACGTCCTATGGTATGCTATGTGGTG CTTAGGGATCCTCTCTTAGAAGAAGATGACAACAAAGACAGCTTCAAGAAAGCAGAGTGA GAATCCCTAGGAGAGATCTTCTTCTACTGTTGTTTCTGTCGAAGTTCTTTCGTCTCACT AGCCTTTCTTGTTCTTCTTCTTCTTCTTATTCATGTGTCTCTCAACCTTTCCAAT TATATTGTTACAAACATTTGCTGTCTAGTTTAAAACATGTAAATGTTTGATGATCTTTGC ATATAACAATGTTTGTAAACGACAGATCAAATTTTGTACATTTACAAACTACTAGAAACG AAGACTCCATTTTTGTTTAAGGTAAACCTTGAATCTCATAGATTGTCGATTGTTGGTATT TTCTGAGGTAAAAACAAATTCCATTTGGAACTTAGAGTATCTAACAGCTAACAACCATAA TCCATTTTCAGGTACGGTTCTGTAGACTGTAGTCTTGCTGACCAGTCCGGCTTAACCACC AGGTAAAAGTCCATGCCAAGACATCTGACATCAGAACGACTGGTCAGGCCGAATTGGTGG ${\tt GGTTTAAAGTTTCTAGAGTgGTTAGTTTTACGACCGACCGGGGTTATATATCTACCCGGT}$

GTTAATCCGTCTAGCTTTACTCTTTAGACCTACCTTAGACAGTTAGACACCTGCTAATTA CAATTAGGCAGATCGAAATGAGAAATCTGGATGGAATCTGTCAATCTGTGGACGATTAAT

Matter No.: 11696-069002 Page 14 of 18 Applicant(s): Sunghwa Choe et al. **DWF 7 MUTANTS** 1250 1230 1210 ATGAGTTTCCTTTTTCTTGTTCAGCAAGTTACCTGTGTTACTTGAGAGTTGAGTTAATGG TACTCAAAGGAAAAAGAACAAGTCGTTCAATGGACACAATGAACTCTCAACTCAATTACC 1270 1310 1290 TAGTAAACGCAATTTAACCCTTATAAGTTTAATCGTATTCAACGAATGACCCAGAGACTT ATCATTTGCGTTAAATTGGGAATATTCAAATTAGCATAAGTTGCTTACTGGGTCTCTGAA 1370 1350 TAAATAAATCCATCGTAACCCTCCACTTCAAAATTCTTTTTAAAAAGTAGCAAATCATTT ATTTATTTAGGTAGCATTGGGAGGTGAAGTTTTAAGAAAAATTTTTCATCGTTTAGTAAA 1410 1390 AAATATTGTAAGTTTGCTTCATTCGAAATTGTAGCTACAGATCTCAAAGCTCCTCCTGTT TTTATAACATTCAAACGAAGTAAGCTTTAACATCGATGTCTAGAGTTTCGAGGAGGACAA 1490 1470 1450 GGCCATATCTCTCTCTAACAAACGCATAGTAACACTTGACCACAGTTTGACTTCTCGGCG CCGGTATAGAGAGAGTTGTTTGCGTATCATTGTGAACTGGTGTCAAACTGAAGAGCCGC 1550 1530 1510 GTTTCATGGCGGCGACTATGGCAGATTATAATGATCAGATCGTCAATGAGACCTCTTTTT CAAAGTACCGCCGCTGATACCGTCTAATATTACTAGTCTAGCAGTTACTCTGGAGAAAAA MAATMADYNDQIVNETSFY 1590 1610 1570 ACAACCGAAtGGTTCTGAGTCACCTTTTGCCGGTGAATCTATGGGAACCTTTACCaCATT TGTTGGCTTaCCAAGACTCAGTGGAAAACGGCcACTTAGATACCCTTGGAAATGGtGTAA N R M V L S H L L P V N L W E P L P H F 1630 1650 1670 TCCTCCAGACATGGCTCCGGAACTACCTCGCCGGAAACATACTCTACTTCATCTCCGGCT AGGAGGTCTGTACCGAGGCCTTGATGGAGCGGCCTTTGTATGAGATGAAGTAGAGGCCGA LQTWLRNYLAGNILYFISGF 1710 1730 1690 TCCTCTGGTGCTTCTACATCTATTACCTTAAACTCAACGTTTACGTCCCCAAAGGTTACT

LWCFYIYYLKLNVYVPK

AGGAGACCACGAAGATGTAGATAATGGAATTTGAGTTGCAAATGCAGGGGTTTCCAATGA

Applicant(s): Sunghwa Choe et al. **DWF 7 MUTANTS** 1790 1750 1770 TTTTTCAATTTCGATGTTCTGTTTTGAAACCTTTCTTTTGTTGATTCCTTCGATTGTATC AAAAAGTTAAAGCTACAAGACAAAACTTTGGAAAGAAAACAACTAAGGAAGCTAACATAG 1850 1830 1810 CGGACTATCTAACACAATATGCAATTGGAAAAAAAGAATGACAATGAAAGTCAAGAACAG 1910 1890 1870 TTCTACTTCTCATTTAATTAGTTTTAAAGTTTAATATTTTTTGGCTAATCCACATTTTTTA AAGATGAAGAGTAAATTAATCAAAA1'TTCAAATTATAAAAACCGATTAGGTGTAAAAAAT 1970 1930 1950 AGTTGAATCTTCCATGAAATTTGAGCTCAAAATATACCATGAAATTGAAATTTGTGGTTC TCAACTTAGAAGGTACTTTAAACTCGAGTTTTATATGGTACTTTAACTTTAAACACCAAG 2010 1990 TTAGTTCTATTTCTTGCTTGGTTTCTTCTATTTTTGTGGTTAGAATCCATTCCTACGAGA AATCAAGATAAAGAACGAACCAAAGAAGATAAAAACACCAATCTTAGGTAAGGATGCTCT ESIPTR 2050 2070 2090 ${\tt AAGGCAATGCTTTTGCAAATATACGTGGCAATGAAGGCTATGCCTTGGTACACTCTTCTT}$ TTCCGTTACGAAAACGTTTATATGCACCGTTACTTCCGATACGGAACCATGTGAGAAGAA KAMLLQIYVAMKAMPWYTLL 2130 2150 2110 CCAGCTGTCTCTGAGTATATGATCGAGCATGGTTGGACCAAATGTTACTCTACACTTGAC GGTCGACAGAGACTCATATACTAGCTCGTACCAACCTGGTTTACAATGAGATGTGAACTG PAVSEYMIEHGWTKCYSTLD 2210 2190 2170 CATTTCAACTGGTTCCTCTGTTTCCTCTACATAGCTCTCTATCTTGTTTTAGTTGAGTTt GTAAAGTTGACCAAGGAGACAAAGGAGATGTATCGAGAGATAGAACAAAATCAACTCAAa H F N W F L C F L Y I A L Y L V L F 2250 2270 2230 ATGATTTATTGGGTTCACAAAGAGCTTCATGACATTAAATTTCTCTATAAGCATCTCCAT TACTAAATAACCCAAGTGTTTCTCGAAGTACTGTAATTTAAAGAGATATTCGTAGAGGTA

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2330 2290 2310 GCTACCCATCATATGTACAACAAGCAAAACACACTCTCTCCATTTGCCGGTATGTCAAAG $\tt CGATGGGTAGTATACATGTTGTTTTGTTGTGAGAGAGGTAAACGGCCATACAGTTTC$ ATHHMYNKONTLSPFA 2390 2350 2370 CTATATGTTCTCAATCTAAATTCAAGAGCTTGTATCAATGGTGACTTCTTTACTTGATGT GATATACAAGAGTTAGATTTAAGTTCTCGAACATAGTTACCACTGAAGAAATGAACTACA 2430 2410 TTTTCGGGTTTTCAGGGCTCGCATTCCATCCGCTGGACGGGATACTTCAGGCTATACCGC AAAAGCCCAAAAGTCCCGAGCGTAAGGTAGGCGACCTGCCCTATGAAGTCCGATATGGCG G L A F H P L D G I L Q A I P H 2490 2510 2470 ACGTGATAGCGCTGTTTATAGTGCCGATTCATCTCATAACACATCTGAGTCTTTTGTTTT TGCACTATCGCGACAAATATCACGGCTAAGTAGAGTATTGTGTAGACTCAGAAAACAAAA VIALFIVPIHLITHLSLLFL 2550 2570 2530 TGGAAGGGATATGGACAGCAAGCATCCATGATTGCATACATGGLAACATCTGGCCTATAA ACCTTCCCTATACCTGTCGTTCGTAGGTACTAACGTATGTACCaTTGTAGACCGGATATT EGIWTASIHDCIHGNIWPIM 2610 2630 2590 TGGGTGCAGGATACCATACCATACACCATACAACATACAAGCATAACTATGGTCATTATA ACCCACGTCCTATGGTATGGTATGTGGTATGTTGTATGTTCGTATTGATACCAGTAATAt G A G Y H T I H H T T Y K H N Y G H Y T 2650 2670 CCATATGGATGGACTGGATGTTTGGCTCTCTTATGGTTCCTTTAGCAGAAAAAGACAGTT GGTATACCTACCtGACCTACAAACCGAGAAATACCAAGGAAATCGTCTTTTTCTGTCAA I W M D W M F G S L M V P L A E K D S F 2730 2750 2710 TCAAGGAGAAAGAAAGTGAGAATGTTCAATGCTCACATGTATTCTTCATATGTTGCTCT AGTTCCTCTTTCTTTTCACTCTTACAAGTTACGAGTGTACATAAGAAGTATACAACGAGA KEKEK* 2770 2790 2810 TCTCGTGACTCTTATTAAAACCTTTCTAATCACTTTGGTGGAATTAAAAACATGACTGCA

FIG. 10E

AGAGCACTGAGAATAATTTTGGAAAGATTAGTGAAACCACCTTAATTTTTGTACTGACGT

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2850

2870

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2890

2830

2910

 ${\tt ATTATATATTGcTGGATGAAGAGTTCAAATTTGGACTAAATCTG}\\ {\tt TAATATTAACGACCTACTTCTCAAGTTTAAACCTGATTTAGAC}$

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1	maatmadynd	givnetsfyn	rmvlshllpv	nlweplphfl	qtwlrnylag
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101	llpavseymi	ehgwtkcyst	ldhfnwflcf	lyialylvlv	efmiywvhke
151	lhdikflykh	lhathhmynk	qntlspfagl	afhpldgilq	aiphvialfi
201	vpihlithls	llflegiwta	sihdcihgni	wpimgagyht	ihhttykhny
251	ghytiwmdwm	fgslmvplae	kdsfkekek		